5



1/6

SEQUENCE LISTING

<110> Gaxiola, Roberto A. Fink, Gerald R. Alper, Seth L.

<120> Proton Transporters And Uses In Plants

<130> 0399.2004-002

<140> US 09/834,998

<141> 2001-04-13

<150> US 09/644,039

<151> 2000-08-22

<150> US 60/164,808

<151> 1999-11-10

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<170> FastSEQ for Windows Version 4.0

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<211> 538

<212> PRT

<213> Artificial Sequence

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<223> Arabidpsis - AtNhx1

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1 5 10 15

His Ala Ser Val Val Ala Leu Asn Leu Phe Val Ala Leu Leu Cys Ala
20 25 30

Cys Ile Val Leu Gly His Leu Leu Glu Glu Asn Arg Trp Met Asn Glu
35 40 45

Ser Ile Thr Ala Leu Leu Ile Gly Leu Gly Thr Gly Val Thr Ile Leu 50 55 60

Leu Ile Ser Lys Gly Lys Ser Ser His Leu Leu Val Phe Ser Glu Asp 70 75 80

Leu Phe Phe Ile Tyr Leu Leu Pro Pro Ile Ile Phe Asn Ala Gly Phe 35 90 95

Gln Val Lys Lys Gln Phe Phe Arg Asn Phe Val Thr Ile Met Leu 100 105 110

Phe Gly Ala Val Gly Thr Ile Ile Ser Cys Thr Ile Ile Ser Leu Gly 115 120 125

Val Thr Gln Phe Phe Lys Lys Leu Asp Ile Gly Thr Phe Asp Leu Gly 130 135 140

Asp Tyr Leu Ala Ile Gly Ala Ile Phe Ala Ala Thr Asp Ser Val Cys 145 150 155 160

Thr Leu Gln Val Leu Asn Gln Asp Glu Thr Pro Leu Leu Tyr Ser Leu

165 170 175 Val Phe Gly Glu Gly Val Val Asn Asp Ala Thr Ser Val Val Val Phe 180 185 190

ORIGINALL CALL

```
Asn Ala Ile Gln Ser Phe Asp Leu Thr His Leu Asn His Glu Ala Ala
                   200
 195
Phe His Leu Leu Gly Asn Phe Leu Tyr Leu Phe Leu Leu Ser Thr Leu
                       215
                                   220
Leu Gly Ala Ala Thr Gly Leu Ile Ser Ala Tyr Val Ile Lys Lys Leu
                   230
                           235
Tyr Phe Gly Arg His Ser Thr Asp Arg Glu Val Ala Leu Met Met Leu
              245
                                 250
Met Ala Tyr Leu Ser Tyr Met Leu Ala Glu Leu Phe Asp Leu Ser Gly
                   265
Ile Leu Thr Val Phe Phe Cys Gly Ile Val Met Ser His Tyr Thr Trp
                           280
His Asn Val Thr Glu Ser Ser Arg Ile Thr Thr Lys His Thr Phe Ala
                       295
                                         300
Thr Leu Ser Phe Leu Ala Glu Thr Phe Ile Phe Leu Tyr Val Gly Met
305
                   310
                                      315
Asp Ala Leu Asp Ile Asp Lys Trp Arg Ser Val Ser Asp Thr Pro Gly
               325
                                  330
Thr Ser Ile Ala Val Ser Ser Ile Leu Met Gly Leu Val Met Val Gly
           340
                              345
Arq Ala Ala Phe Val Phe Pro Leu Ser Phe Leu Ser Asn Leu Ala Lys
                           360
                                              365
Lys Asn Gln Ser Glu Lys Ile Asn Phe Asn Met Gln Val Val Ile Trp
                       375
                                          380
Trp Ser Gly Leu Met Arg Gly Ala Val Ser Met Ala Leu Ala Tyr Asn
                   390
                                      395
Lys Phe Thr Arg Ala Gly His Thr Asp Val Arg Gly Asn Ala Ile Met
              405
                                  410
                                                      415
Ile Thr Ser Thr Ile Thr Val Cys Leu Phe Ser Thr Val Val Phe Gly
           420
                               425
Met Leu Thr Lys Pro Leu Ile Ser Tyr Leu Leu Pro His Gln Asn Ala
                           440
Thr Thr Ser Met Leu Ser Asp Asp Asn Thr Pro Lys Ser Ile His Ile
                       455
Pro Leu Leu Asp Gln Asp Ser Phe Ile Glu Pro Ser Gly Asn His Asn
465
                 470
                                      475
Val Pro Arg Pro Asp Ser Ile Arg Gly Phe Leu Thr Arg Pro Thr Arg
               485
                                  490
Thr Val His Tyr Tyr Trp Arg Gln Phe Asp Asp Ser Phe Met Arg Pro
           500
                              505
Val Phe Gly Gly Arg Gly Phe Val Pro Phe Val Pro Gly Ser Pro Thr
                          520
                                             525
Glu Arg Asn Pro Pro Asp Leu Ser Lys Ala
                       535
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Met Ala Arg Arg Gly Trp Arg Arg Ala Pro Leu Arg Arg Gly Val Gly
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                                 10
Ser Ser Pro Arg Ala Arg Arg Leu Met Arg Pro Leu Trp Leu Leu Leu
           20
                              25
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Ala Val Gly Val Phe Asp Trp Ala Gly Ala Ser Asp Gly Gly Gly Glu Ala Arg Ala Met Asp Glu Glu Ile Val Ser Glu Lys Gln Ala Glu Glu Ser His Arg Gln Asp Ser Ala Asn Leu Leu Ile Phe Ile Leu Leu 7.0 Leu Thr Leu Thr Ile Leu Thr Ile Trp Leu Phe Lys His Arg Arg Ala Arg Phe Leu His Glu Thr Gly Leu Ala Met Ile Tyr Gly Leu Leu Val Gly Leu Val Leu His Tyr Gly Ile His Val Pro Ser Asp Val Asn Asn Val Thr Leu Ser Cys Glu Val Gln Ser Ser Pro Thr Thr Leu Leu Val Thr Phe Asp Pro Glu Val Phe Phe Asn Ile Leu Leu Pro Pro Ile Ile Phe Tyr Ala Gly Tyr Ser Leu Lys Arg Arg His Phe Phe Arg Asn Leu Gly Ser Ile Leu Ala Tyr Ala Phe Leu Gly Thr Ala Ile Ser Cys Phe Val Ile Gly Ser Ile Met Tyr Gly Gly Val Thr Leu Met Lys Val Thr Gly Gln Leu Ala Gly Asp Phe Tyr Phe Thr Asp Cys Leu Leu Phe Gly Ala Ile Val Ser Ala Thr Asp Pro Val Thr Val Leu Ala Ile Phe His Glu Leu Gln Val Asp Val Glu Leu Tyr Ala Leu Leu Phe Gly Glu Ser Val Leu Asn Asp Ala Val Ala Ile Val Leu Ser Ser Ser Ile Val Ala Tyr Gln Pro Ala Gly Asp Asn Ser His Thr Phe Asp Val Thr Ala Met Phe Lys Ser Ile Gly Ile Phe Leu Gly Ile Phe Ser Gly Ser Phe Ala Met Gly Ala Ala Thr Gly Val Val Thr Ala Leu Val Thr Lys Phe Thr Lys Leu Arg Glu Phe Gln Leu Leu Glu Thr Gly Leu Phe Phe Leu Met Ser Trp Ser Thr Phe Leu Leu Ala Glu Ala Trp Gly Phe Thr Gly Val Val Ala Val Leu Phe Cys Gly Ile Thr Gln Ala His Tyr Thr Tyr Asn Asn Leu Ser Thr Glu Ser Gln His Arg Thr Lys Gln Leu Phe Glu Leu Leu Asn Phe Leu Ala Glu Asn Phe Ile Phe Ser Tyr Met Gly Leu Thr Leu Phe Thr Phe Gln Asn His Val Phe Asn Pro Thr Phe Val Val Gly Ala Phe Val Ala Ile Phe Leu Gly Arg Ala Ala Asn Ile Tyr Pro Leu Ser Leu Leu Leu Asn Leu Gly Arg Arg Ser Lys Ile Gly Ser Asn Phe Gln His Met Met Met Phe Ala Gly Leu Arg Gly Ala Met Ala Phe Ala Leu Ala Ile Arg Asp Thr Ala Thr Tyr Ala Arg Gln Met Met Phe Ser Thr Thr Leu Leu Ile Val Phe Phe Thr Val Trp Val Phe Gly Gly Gly

```
Thr Thr Ala Met Leu Ser Cys Leu His Ile Arg Val Gly Val Asp Ser
                               505
Asp Gln Glu His Leu Gly Val Pro Glu Asn Glu Arg Arg Thr Thr Lys
                          520
Ala Glu Ser Ala Trp Leu Phe Arg Met Trp Tyr Asn Phe Asp His Asn
                      535
Tyr Leu Lys Pro Leu Leu Thr His Ser Gly Pro Pro Leu Thr Thr
                  550
                                     555
Leu Pro Ala Cys Cys Gly Pro Ile Ala Arg Cys Leu Thr Ser Pro Gln
              565
                                 570
Ala Tyr Glu Asn Gln Glu Gln Leu Lys Asp Asp Ser Asp Leu Ile
           580
                             585
Leu Asn Asp Gly Asp Ile Ser Leu Thr Tyr Gly Asp Ser Thr Val Asn
 595
                          600
                                             605
Thr Glu Pro Ala Thr Ser Ser Ala Pro Arg Arg Phe Met Gly Asn Ser
                                         620
                      615
Ser Glu Asp Ala Leu Asp Arg Glu Leu Ala Phe Gly Asp His Glu Leu
                  630
                                     635
Val Ile Arg Gly Thr Arg Leu Val Leu Pro Met Asp Asp Ser Glu Pro
              645
                                 650
Pro Leu Asn Leu Leu Asp Asn Thr Arg His Gly Pro Ala
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<210> 3

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<211> 633
<212> PRT
<213> Artificial Sequence
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<223> Yeast - ScNhx1
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                                   10
Thr Ala Lys Arg Ala Val Asp Pro Asp Asp Asp Glu Leu Leu Pro
           20
                               25
Ser Pro Asp Leu Pro Gly Ser Asp Pro Ile Ala Gly Asp Pro Asp
                           40
                                               45
Val Asp Leu Asn Pro Val Thr Glu Glu Met Phe Ser Ser Trp Ala Leu
                       55
Phe Ile Met Leu Leu Leu Ile Ser Ala Leu Trp Ser Ser Tyr Tyr
                   70
                                       75
Leu Thr Gln Lys Arg Ile Arg Ala Val His Glu Thr Val Leu Ser Ile
               85
                                   90
Phe Tyr Gly Met Val Ile Gly Leu Ile Ile Arg Met Ser Pro Gly His
           100
                               105
Tyr Ile Gln Asp Thr Val Thr Phe Asn Ser Ser Tyr Phe Phe Asn Val
       115
                           120
                                               125
Leu Leu Pro Pro Ile Ile Leu Asn Ser Gly Tyr Glu Leu Asn Gln Val
                       135
                                           140
Asn Phe Phe Asn Asn Met Leu Ser Ile Leu Ile Phe Ala Ile Pro Gly
                  150
                                      155
Thr Phe Ile Ser Ala Val Val Ile Gly Ile Ile Leu Tyr Ile Trp Thr
                       170
                                                      175
Phe Leu Gly Leu Glu Ser Ile Asp Ile Ser Phe Ala Asp Ala Met Ser
           180
```

Val	Gly	Ala 195	Thr	Leu	Ser	Ala	Thr	Asp	Pro	Val	Thr	Ile 205	Leu	Ser	Ile
Phe	Asn 210	Ala	Tyr	Lys	Val	Asp 215		L∵s	Leu	Tyr	Thr 220	Ile	Ile	Phe	Gly
Glu 225	Ser	Leu	Leu	Asn	Asp 230	Ala	Ile	Ser	Ile	Val 235	Met	Phe	Glu	Thr	Cys 240
Gln	Lys	Phe	His	Gly 245	Gln	Pro	Ala	Thr	Phe 250	Ser	Ser	Val	Phe	Glu 255	Gly
Ala	Gly	Leu	Phe 260	Leu	Met	Thr	Phe	Ser 265	Val	Ser	Leu	Leu	Ile 270	Gly	Val
Leu	Ile	Gly 275	Ile	Leu	Val	Ala	Leu 280	Leu	Leu	Lys	His	Thr 285	His	Ile	Arg
Arg	Tyr 290	Pro	Gln	Ile	Glu	Ser 295	C','s	Leu	Ile	Leu	Leu 300	Ile	Ala	Tyr	Glu
Ser 305	Tyr	Phe	Phe	Ser	Asn 310	Gly	C∵s	His	Met	Ser 315	Glγ	Ile	Val	Ser	Leu 320
				325					330					Met 335	
			340					345					350	Ala	
		355					360					365		Phe	
	370				-	375					380			Ala	
385					390					395				Gln	400
		_		405	_				410					Gly 415	
	_		420					425					430	Tyr	
		435		_			440					445		Ala	
	450	_			_	455		_			460			Thr	
465					470					475	_	_		Thr	480
_				485					490					Glu 495	
			500					505					510	Ile	
		515					510					525		Ser	
	530					535					540			Ser	
545					550					555				Phe	560
_				565					570					Ser 575	
	_	_	580					585					590	Asn -	
_		595	_				6 C O					605	_	Pro	
	61)	-				615			GIn	Asp	Ser 620	Ala	Thr	Gln	ser
Pro 625	Ala	Asp	rne	ser	3er 630	Gin	Asn	HIS							

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<213> Unknown
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Thr Cys Thr Ala Gly Thr Gly Thr Cys Gly Ala Ala Ala Cys Thr Gly
            20
                                25
Cys Cys Thr Thr Cys Gly
       35
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